

# Notation

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Let  $K > 0 \in \mathbb{R}$  be a caliper. Then, let schools be indexed  $i = 1, \dots, S$  and for each school  $i$ , students be indexed  $j = 1, \dots, N_i$ . Let  $N = \sum_{i=1}^S N_i$ . Treatment is assigned at the school-level, so let  $T_i \in \{0, 1\}$  indicate whether a school received treatment or not. Then, let  $Y_{ij} \in \mathbb{R}$  be the outcome for student  $(i, j)$  and  $\hat{Y}_{ij} \in \mathbb{R}$  be the estimated prognostic score for student  $(i, j)$ . Let  $T \subset [S]$  be the set of treatment schools and  $C = [S] \setminus T$  be the set of control schools. Let  $S_{ij}$  be the school that student  $(i, j)$  attends. All future data products are functions of these objects.

There are four final data products:

1.  $(\ell_M, u_M)$ : the confidence interval from the **treatment effect estimation** derived using the **matchAhead procedure**.
2.  $(\ell_P, u_P)$ : the confidence interval from the **treatment effect estimation** derived using the **Pimentel procedure**.
3.  $\{E_{ij}^{(M)}\}_{T \times C}$  the set of times elapsed when matching treatment to control schools using the using the **matchAhead procedure**.
4.  $\{E_{ij}^{(P)}\}_{T \times C}$  the set of times elapsed when matching treatment to control schools using the using the **Pimentel procedure**.